Amendments

This listing of claims will replace all prior versions, and listings of claims in the application.

- (Currently amended) An isolated bacterium comprising: an F episome that confers high efficiency transformability; one or more mutations that allow transformation of methylated nucleic acids; one or <u>more</u> mutations that allow transformation with unmethylated nucleic acids; and one or more mutations that confer resistance to bacteriophage infection.
- (Original) The bacterium of claim 1 wherein said bacterium is an Escherichia coli strain.
 - 3. (Original) The bacterium of claim 2 wherein said strain is K-12.
- 4. (Original) The bacterium of claim 3 wherein the genotype of said F episome comprises $proAB^{+}$ lacf q $lacZ\Delta M15$ Tn10(Tet^{R}).
- (Original) The bacterium of claim 3 wherein said one or more mutations that allow transformation of unmethylated nucleic acids involve the hsdR, hsdM and/or hsdS genes.
- (Original) The bacterium of claim 3 wherein said one or more mutations that allow transformation of methylated nucleic acids involve the mcrA, mcrBC and/or mrr genes.
 - 7. (Original) The bacterium of claim 3 wherein said bacteriophage is phage T1.
- 8. (Original) The bacterium of claim 3 wherein said one or more mutations that confer resistance to bacteriophage infection involve the tonA gene.

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- (Original) An isolated Escherichia coli K-12 bacterium, said bacterium having a genotype comprising mcrA Δ(mrr-hsdRMS-mcrBC) tonA / F^{*} proAB^{*} lacf[§] lacZΔMS15 Tn10(Tet^B).
- 10. (Original) The bacterium of claim 9, wherein said genotype comprises: mcrA Δ(mrr-hsdRMS-mcrBC) Φ80(lacZ)ΔM15 Δ(lacZYA-argF) U169 endΔ1 recΔ1 supE44 thi-1 gyrΔ96 relΔ1 deoR tonΔ panD / F proΔB* lacI^q lacZΔM15 Tn10(Tet^R).
- (Original) The bacterium of claim 9, wherein said bacterium is Escherichia coli BRL3946 (NRRL accession No. B-30640).
- (Original) The bacterium of claim 9, wherein said bacterium is a derivative of Escherichia coli BRL3946 (NRRL accession No. B-30640).